OIPE

Page 1 of 7

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/001,227

DATE: 12/10/2001 TIME: 15:00:38

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\12102001\1001227.raw



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              SILOS-SANTIAGO
      7 <120> TITLE OF INVENTION: METHODS OF USING 18903 TO TREAT PAIN AND
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     10 <130> FILE REFERENCE: MNI-199
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/001,227
C--> 12 <141> CURRENT FILING DATE: 2001-11-30
     12 <150> PRIOR APPLICATION NUMBER: 60/250929
     13 <151> PRIOR FILING DATE: 2000-11-30
     15 <160> NUMBER OF SEQ ID NOS: 7
     17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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    26 <222> LOCATION: (98)...(1849)
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RAW SEQUENCE LISTING DATE: 12/10/2001 PATENT APPLICATION: US/10/001,227 TIME: 15:00:38

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\12102001\I001227.raw

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74 ttc atc gtg ggc gct tct tcg tac gag ggc tct gac ttg gcc gcc 75 Phe Ile Val Gly Ala Ala Ser Ser Tyr Glu Gly Ser Asp Leu Ala Ala 76 170 180 175 180 180 175 180 175 180 175 180 175 180 175 180 175 180 175 180 175 180 175 180 175 180 175 180 175 180 175 180 175 180 175 180 180 185 190 195 195 195 195 195 195 195 195 195 195	72	110	OI,	изр	rio	155	ьеu	PIO	Val	мес	160	Trp	Phe	Pro	Gly		Ala	
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86 ctg gac cag atg gcg gct ctg cgc tgg gtg cag gag aac atc gca gcc 87 Reu Asp Gln Met Ala Ala Leu Arg Trp Val Gln Glu Asn Ile Ala Ala 225 220 220 225 230 230 250 260 1 Phe Gly Gly Asp Pro Gly Asn Val Thr Leu Phe Gly Gln Ser Ala Gly 235 240 245 245 245 240 245 245 245 245 245 245 245 245 245 245	84	_	200				p	205	JCI	птэ	АІа	Arg		ASI	Trp	GLY	Leu	
88 215	86	ctg	gac	cag	atg	gcg	gct	ctg	cgc	tgg	qtq	caq	σασ	aac	atc	aca	acc	787
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92 235 240 245 88. 94 gcc atg agc atc tca gga ctg atg atg tca ccc cta gcc tcg ggt ctc 95 Ala Met Ser Ile Ser Gly Leu Met Met Ser Pro Leu Ala Ser Gly Leu 96 250 260 98 ttc cat cgg gcc att tcc cag agt ggc acc gcg tta ttc aga ctt ttc 99 Phe His Arg Ala Ile Ser Gln Ser Gly Thr Ala Leu Phe Arg Leu Phe 100 265 270 102 atc act agt aac cca ctg aaa gtg gcc aag aag gtt gcc cac ctg gct 103 Ile Thr Ser Asn Pro Leu Lys Val Ala Lys Lys Val Ala His Leu Ala 104 280 285 106 gga tgc aac cac aac ag aca cag atc ctg gta aac tgc ctg agg gca 107 Gly Cys Asn His Asn Ser Thr Gln Ile Leu Val Asn Cys Leu Arg Ala 108 295 300 305 310 110 cta tca ggg acc aag gtg atg cgt gtg tcc aac aag atg aga ttc ctc 111 Leu Ser Gly Thr Lys Val Met Arg Val Ser Asn Lys Met Arg Phe Leu 112 315 320 325 114 caa ctg aac ttc cag aga gac ccg gaa gag att atc tgg tcc atg agc 115 Gln Leu Asn Phe Gln Arg Asp Pro Glu Glu Ile Ile Trp Ser Met Ser 116 330 335 340 118 cct gtg gtg gat ggt gtg gtg atc cca gat gac cct ttg gtg ctc ctc 119 Pro Val Val Asp Gly Val Val Ile Pro Asp Asp Pro Leu Val Asn Asn 124 360 365 370 125 Leu Glu Phe Asn Trp Leu Leu Cct tat atc atg aag ttc ccg cta aac 125 Leu Glu Phe Asn Trp Leu Leu Phe Asp Coc Cac cta gcc tac ctt cac gcd cac acc 126 Leu Glu Phe Asn Trp Leu Leu Phe 127 Leu Glu Phe Asn Trp Leu Leu Cct tat atc atg aag ttc ccg cta aac 128 375 380	00	213					220					225					230	
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108 295 300 305 310 110 cta tca ggg acc aag gtg atg cgt gtg tcc aac aag atg aga ttc ctc 111 Leu Ser Gly Thr Lys Val Met Arg Val Ser Asn Lys Met Arg Phe Leu 112 315 320 325 114 caa ctg aac ttc cag aga gac ccg gaa gag att atc tgg tcc atg agc 115 Gln Leu Asn Phe Gln Arg Asp Pro Glu Glu Ile Ile Trp Ser Met Ser 116 330 335 340 118 cct gtg gtg gat ggt gtg gtg atc cca gat gac cct ttg gtg ctc ctg 119 Pro Val Val Asp Gly Val Val Ile Pro Asp Asp Pro Leu Val Leu Leu 120 345 350 355 122 acc cag ggg aag gtt tca tct gtg ccc tac ctt cta ggt gtc aac aac 123 Thr Gln Gly Lys Val Ser Ser Val Pro Tyr Leu Leu Gly Val Asn Asn 124 360 365 370 126 ctg gaa ttc aat tgg ctc ttg cct tat atc atg aag ttc ccg cta aac 127 Leu Glu Phe Asn Trp Leu Leu Pro Tyr Ile Met Lys Phe Pro Leu Asn 128 375 380 385	107	СТА	Cys	Asn	His	Asn	Ser	Thr	Gln	Ile	Leu	Val	Asn	Cys	Leu	Arg	Ala	1027
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112 315 320 325 114 caa ctg aac ttc cag aga gac ccg gaa gag att atc tgg tcc atg agc 11 115 Gln Leu Asn Phe Gln Arg Asp Pro Glu Glu Ile Ile Trp Ser Met Ser 116 330 335 340 118 cct gtg gtg gat ggt gtg gtg atc cca gat gac cct ttg gtg ctc ctg 11 119 Pro Val Val Asp Gly Val Val Ile Pro Asp Asp Pro Leu Val Leu Leu 20 345 350 355 122 acc cag ggg aag gtt tca tct gtg ccc tac ctt cta ggt gtc aac aac 12 123 Thr Gln Gly Lys Val Ser Ser Val Pro Tyr Leu Leu Gly Val Asn Asn 260 365 370 126 ctg gaa ttc aat tgg ctc ttg cct tat atc atg aag ttc ccg cta aac 12 127 Leu Glu Phe Asn Trp Leu Leu Pro Tyr Ile Met Lys Phe Pro Leu Asn 380 385	111	CLd Lou	Cor	ggg	acc	aag	gtg	atg	cgt	gtg	tcc	aac	aag	atg	aga	ttc	ctc	1075
114 caa ctg aac ttc cag aga gac ccg gaa gag att atc tgg tcc atg agc 11 115 Gln Leu Asn Phe Gln Arg Asp Pro Glu Glu Ile Ile Trp Ser Met Ser 330 335 340 118 cct gtg gtg gat ggt gtg gtg atc cca gat gac cct ttg gtg ctc ctg 11 119 Pro Val Val Asp Gly Val Val Ile Pro Asp Asp Pro Leu Val Leu Leu 120 345 350 355 122 acc cag ggg aag gtt tca tct gtg ccc tac ctt cta ggt gtc aac aac 12 123 Thr Gln Gly Lys Val Ser Ser Val Pro Tyr Leu Leu Gly Val Asn Asn 124 360 365 370 126 ctg gaa ttc aat tgg ctc ttg cct tat atc atg aag ttc ccg cta aac 12 127 Leu Glu Phe Asn Trp Leu Leu Pro Tyr Ile Met Lys Phe Pro Leu Asn 128 375 380 385	112	Leu	261	СТУ	1111	цуS 315	Val	мет	Arg	Val	Ser	Asn	Lys	Met	Arg		Leu	
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122 acc cag ggg aag gtt tca tct gtg ccc tac ctt cta ggt gtc aac aac 12 123 Thr Gln Gly Lys Val Ser Ser Val Pro Tyr Leu Leu Gly Val Asn Asn 124 360 365 370 126 ctg gaa ttc aat tgg ctc ttg cct tat atc atg aag ttc ccg cta aac 12 127 Leu Glu Phe Asn Trp Leu Leu Pro Tyr Ile Met Lys Phe Pro Leu Asn 128 375 380 385	TIJ	Pro	Val	val	Asp	Gly	Val	Val	Ile	Pro	Asp	Asp	Pro	Leu	Val	Leu	Leu	
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124 360 365 370 126 ctg gaa ttc aat tgg ctc ttg cct tat atc atg aag ttc ccg cta aac 12 127 Leu Glu Phe Asn Trp Leu Leu Pro Tyr Ile Met Lys Phe Pro Leu Asn 128 375 380 385	123	Thr	Gln	999 Gl v	Lvs	Val	Ser	Sor	gtg	CCC	tac	ctt	cta	ggt	gtc	aac	aac	1219
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RAW SEQUENCE LISTING

DATE: 12/10/2001 PATENT APPLICATION: US/10/001,227 TIME: 15:00:38

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\12102001\1001227.raw

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135 Arg Thr Leu Leu Asn Ile Thr Lys Glu Gln Val Pro Leu Val Val Glu 410	
138 gag tac ctg gac aat gtc aat gag cat gac tgg aag atg cta cga aac 14: 139 Glu Tyr Leu Asp Asn Val Asn Glu His Asp Trp Lys Met Leu Arg Asn	11
142 cgt atg atg gac ata gtt caa gat ggg act the att	
143 Arg Met Met Asp Ile Val Gln Asp Ala Thr Phe Val Tyr Ala Thr Leu	59
445	
146 cag act gct cac tac cac cga gat ggg ggg ats and the	
of the first are Ard ASD Ala GIV Len Dro Val Turn Law Terr)7
400 /65	
150 gaa ttt gag cac cac gct cgt gga ata atg gtg ana	
of the old his his Ald Ard GIV [[e T]e Val Ive Dwo Ame with	, 5
4/3 /80	
134 999 yea gae cat gag gat gag atg tag tto gto tto	
and the his diy Asp Giu Met Tyr Phe Leu Phe Gly Gly Pro Phe	3
450	
130 gcc aca ggc ctt tcc atg ggt aag gag aag aag	1
160 Leu Ser Met Gly Lys Glu Lys Ala Leu Ser Leu Gln Met	_
7 515	
162 atg aaa tac tgg gcc aac ttt gcc cgc aca gga aac ccc aat gat ggg 169	9
164 520 FOR	
37.) 530	
166 aat ctg ccc tgc tgg cca cgc tac aac aag gat gaa aag tac ctg cag 174'	7
167 Asn Leu Pro Cys Trp Pro Arg Tyr Asn Lys Asp Glu Lys Tyr Leu Gln 168 535 540 545	
170 ctg gat ttt acc aca aga gtg ggc atg aag ctc aag gag aag aag atg 1795	
171 Leu Asp Phe Thr Thr Arg Val Gly Met Lys Leu Lys Glu Lys Lys Met	5
555	
1/4 gct ttt tgg atg agt ctg tac cag tgt gap agg	
The rip Met Ser Leu Tyr Gin Ser Gin Arg Dro Clu Tue Gin	}
5/5	
1/0 Cad LLC taagggtggc tatgcaggaa ggaggcaaag aggggttt	
110	
182 caggecetgg ggagactage catggacata cetggggaca agagttetae ecaagggega 1959	,
TO THE SECTION OF THE	
186 <211> LENGTH: 584	
187 <212> TYPE: PRT	
188 <213> ORGANISM: Homo sapiens 190 <400> SEQUENCE: 2	
191 Met Dro Com Mbm Wall to -	
191 Met Pro Ser Thr Val Leu Pro Ser Thr Val Leu Pro Ser Leu Leu Pro	
193 Thr Ala Gly Ala Gly Trp Ser Met Arg Trp Ile Leu Cys Trp Ser Leu 25	
195 Thr Leu Cys Leu Met Ala Gln Thr Ala Leu Gly Ala Leu His Thr Lys	

RAW SEQUENCE LISTING

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Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\12102001\I001227.raw

196	:		2 5													
		~ D	35					40					45	,		
		50					ככ					60				n Met
199	Hi	s Va	1 G1	y Lv	s Th	r Pr	o Il	e G1:	n Va	1 ph	O T O		17-	1 5	_,	e Ser
200	65					70	- 11	C 01.	• •	T F1	75	u GI	y va	T Pr	o Ph	
201	Arc	a Pro	o Pro	o Lei	1 G1	v T1.	a T.ai	1 7 200	a Dh		75					80 o Glu
202		,		J	85	y	е пе	u Ar	y Pii	e Al	a Pr	o Pr	o GI	u Pr		
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204		1	Y LIY	100	J А. ТТ (S AL	y ASI	O AL	a Th	r Th -	r Ty	r Pr	o Pr	o G1	у Су	s Leu
		C1.	1 60.			- 01	_		10	5				11	0	
206	GII	ı GI	1 50.	r Trg	GT2	y Gli	n Lei	ı Ala	a Se	r Me	t Ty:	r Va	l Se	r Th	r Ar	g Glu
200			44.	,				120)				1 2	5		
207	Arg	туг	с гуз	s Trp) Let	ı Arg	J Ph∈	e Sei	Gl:	ı As	р Су	s Le	u Ty	r Lei	ı As	n Val
		100	,				135)				1/	Λ			
209	Tyr	` Ala	Pro) Ala	ı Arç	y Ala	a Pro	Gly	Ası	Pr	o Gli	ı Lei	ı Pro	o Vai	l Me	t Val
210	T 4 7	,				100)				15	ξ				1 60
211	Trp	Phe	Pro	Gly	Gly	7 Ala	≀ Ph∈	: Ile	· Vai	L G1	v Ala	1 Δ1:	3 Co.	r Soi	^ M++	160 r Glu
212					T 0 3)				174	Λ				1 7	_
213	Gly	Ser	Asp	Leu	Ala	Ala	. Ara	Glu	T.v.s	Va.	0 1 1751	T O	. 170	l Dl.	17	ı Gln
214			-	180			9	010	185	, vu.	T 401	. ье	ı va.			ı Gln
215	His	Ara	Leu	Glv	Tle	Dhe	. G1v	Dho	TOU	,	. mb		_	190) .	s Ala
216		,	195			. 1 110	GLY	200	пес	ı sei	r Thr	ASE	Asp	Ser	His	s Ala
217	Ara	Glv			C117	. T 011	Tou	200	a 1.			_	205	5		
218	9	210	Hon	тър	СТУ	ьeu	Leu	Asp	GIR	Met	: Ala	Ala	Let	ı Arg	Tr	Val
	Gln			т1.			215					220)			
220	335	GIU	ASII	тте	Ата	Ala	Phe	Gly	Gly	Asp	Pro	Gly	Asr	val	Thi	Leu
	423					230					225					0.40
221	Pne	GTA	GIn	Ser	Ala	Gly	Ala	Met	Ser	Ile	e Ser	Gly	Leu	Met	Met	240 Ser
222					245					250	1				255	
223	Pro	Leu	Ala	Ser	Gly	Leu	Phe	His	Arg	Ala	Ile	Ser	Gln	Ser	Glv	Thr
				200					265					270		
225	Ala	Leu	Phe	Arg	Leu	Phe	Ile	Thr	Ser	Asn	Pro	Len	T.ve	Val	7.1.5	Trra
			2/3					280					205			
227	Lys	Val	Ala	His	Leu	Ala	Glv	Cvs	Asn	Hic	Asn	Cor	mb~	01 -	- 1 -	_
228		290					295	-1-		1110	11511	300	1111	GIII	тте	Leu
229	Val	Asn	Cvs	Leu	Ara	Ala	Len	Sor	C117	m h m	T	300		_		
230	305		1 -		5	310	пси	Ser	GTĀ	T 111T	Lys	val	met	Arg	Val	Ser
		Lvs	Met	Ara	Dho	LAU	Cln	T 011	7 ~ ~	Dl.	315	_				320
231 z 232		-1-		9	325	пец	GIII	ьеu	ASII	Pne	GIn	Arg	Asp	Pro	Glu	Glu
	Tle	Tla	Trn	Cor		O	D			330	_				335	
233 : 234		110	тъ	240	мес	ser	Pro	val	Val	Asp	Gly	Val	Val	Ile	Pro	Asp
	۱an	Dro	т о	340	- -	_			345					350		
235 Z 236	asp	PIO	Leu	val	ьeu	Leu	Thr	Gln	Gly	Lys	Val	Ser	Ser	Val	Pro	Tyr
•			J J J					100					265			
237 I 238	Leu	Leu	Gly	Val	Asn	Asn	Leu	Glu	Phe	Asn	Trp	Leu	Leu	Pro	Tvr	Tle
		5,0					3/3					3 ይሰ				
239 N	1et	Lys	Phe	Pro	Leu	Asn	Arg	Gln	Ala	Met	Ara	Lvs	Glu	Thr	Tla	Thr
	,05					390					395					400
241 I 242	ys :	Met	Leu	Trp	Ser	Thr	Ara	Thr	Leu	Len	Agn	Tla	ጥሎ∽	T 17.0	c1	~± U U
					403					410					41-	
243 V	al :	Pro :	Leu	Val	Val	Glu	Glu	ηνr	Τ.Δ11	170	λαν	17 n 1	7 ~	~ 1	415	_
244				420			J_ u	-1-	425	uah	ASII	val	ASN		HlS	Asp
				•					4 4 J					430		

RAW SEQUENCE LISTING DATE: 12/10/2001 PATENT APPLICATION: US/10/001,227 TIME: 15:00:38

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\12102001\I001227.raw

```
245 Trp Lys Met Leu Arg Asn Arg Met Met Asp Ile Val Gln Asp Ala Thr
                                  440
 247 Phe Val Tyr Ala Thr Leu Gln Thr Ala His Tyr His Arg Asp Ala Gly
 248
                              455
 249 Leu Pro Val Tyr Leu Tyr Glu Phe Glu His His Ala Arg Gly Ile Ile
                          470
                                               475
 251 Val Lys Pro Arg Thr Asp Gly Ala Asp His Gly Asp Glu Met Tyr Phe
                      485
                                           490
 253 Leu Phe Gly Gly Pro Phe Ala Thr Gly Leu Ser Met Gly Lys Glu Lys
                  500
                                      505
 255 Ala Leu Ser Leu Gln Met Met Lys Tyr Trp Ala Asn Phe Ala Arg Thr
             515
                                  520
 257 Gly Asn Pro Asn Asp Gly Asn Leu Pro Cys Trp Pro Arg Tyr Asn Lys
 258
         530
                              535
 259 Asp Glu Lys Tyr Leu Gln Leu Asp Phe Thr Thr Arg Val Gly Met Lys
 260 545
                          550
                                              555
 261 Leu Lys Glu Lys Lys Met Ala Phe Trp Met Ser Leu Tyr Gln Ser Gln
                     565
                                          570
 263 Arg Pro Glu Lys Gln Arg Gln Phe
 264
                 580
 267 <210> SEQ ID NO: 3
 268 <211> LENGTH: 1752
 269 <212> TYPE: DNA
 270 <213> ORGANISM: Homo sapiens
 272 <220> FEATURE:
 273 <221> NAME/KEY: CDS
 274 <222> LOCATION: (1)...(1752)
 276 <400> SEQUENCE: 3
277 atg cca tcc aca gtg ttg cca tcc aca gtg ttg cca tca ctc ctg ccc
                                                                        48
278 Met Pro Ser Thr Val Leu Pro Ser Thr Val Leu Pro Ser Leu Leu Pro
279 1
281 aca gca gga gct ggc tgg agc atg agg tgg att ctg tgc tgg agc ctc
                                                                        96
282 Thr Ala Gly Ala Gly Trp Ser Met Arg Trp Ile Leu Cys Trp Ser Leu
                                      25
285 acc ctc tgc ctg atg gcg cag acg gcc ttg ggt gcc ttg cac acc aag
                                                                        144
286 Thr Leu Cys Leu Met Ala Gln Thr Ala Leu Gly Ala Leu His Thr Lys
287
             35
                                  40
289 agg cct caa gtg gtc acc aaa tat gga acc ctg caa gga aaa cag atg
290 Arg Pro Gln Val Val Thr Lys Tyr Gly Thr Leu Gln Gly Lys Gln Met
                              55
293 cat gtg ggg aag aca ccc atc caa gtc ttt tta gga gtc ccc ttc tcc
                                                                        240
294 His Val Gly Lys Thr Pro Ile Gln Val Phe Leu Gly Val Pro Phe Ser
                         70
297 aga cct cct cta ggt atc ctc agg ttt gca cct cca gaa ccc ccg gag
                                                                        288
298 Arg Pro Pro Leu Gly Ile Leu Arg Phe Ala Pro Pro Glu Pro Pro Glu
                     85
301 ccc tgg aaa gga atc aga gat gct acc acc tac ccg cct ggg tgc ctg
302 Pro Trp Lys Gly Ile Arg Asp Ala Thr Thr Tyr Pro Pro Gly Cys Leu
303
                100
                                     105
                                                         110
```



Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/001,227

DATE: 12/10/2001 TIME: 15:00:39

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\12102001\I001227.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 L:494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5